IN THE SPECIFICATION

Kindly amend the specification by replacement of the following.

Page 6, replace the paragraph starting at line 28 with:

The peptide epitope is typically a fragment of any of the pathogen proteins mentioned. In the case of M. tuberculosis the peptide epitope may be any of the peptides shown below from ESAT-6 (SEQ ID NO: 36) and CFP-10 (SEQ ID NO: 37).

Page 6, replace the list starting at line 32 with:

Peptides from ESAT-6 (SEQ ID NOS: 1	17, respectively):
MTEQQWNFAGIEAAA	(SEQ. ID NO: 1)
WNFAGIEAAASAIQG	(SEQ. ID NO: 2)
IEAAASAIQGNVTSI	(SEQ. ID NO: 3)
SAIQGNVTSIHSLLD	(SEQ. ID. NO: 4)
NVTSIHSLLDEGKQS	(SEQ. ID. NO: 5)
HSLLDEGKQSLTKLA	(SEQ. ID NO: 6)
EGKQSLTKLAAAWGG	(SEQ. ID.NO: 7)
LTKLAAAWGGSGSEA	(SEQ. ID.NO: 8)
AAWGGSGSEAYQGVQ	(SEQ. ID. NO: 9)
SGSEAYQGVQQKWDA	(SEQ. ID. NO: 10)
YQGVQQKWDATATEL	(SEQ. ID. NO: 11)
QKWDATATELNNALQ	(SEQ. ID. NO: 12)
TATELNNALQNLART	(SEQ. ID. NO: 13)
NNALQNLARTISEAG	(SEQ. ID. NO: 14)
NLARTISEAGQAMAS	(SEQ. ID. NO: 15)
ISEAGQAMASTEGNV	(SEQ. ID. NO: 16)

Page 7, replace the list starting at line 20 with:

Peptides from CFP-10 (SEQ ID NOS: 18 - 35, respectively):

M A E M K T D A A T L A Q E A (SEQ ID NO: 18)

QAMASTEGNVTGMFA

(SEQ. ID NO: 17)

TDAATLAQEAGNFER	(SEQ ID NO: 19)
LAQEAGNFERISGDL	(SEQ ID NO: 20)
GNFERISGDLKTQID	(SEQ ID NO: 21)
ISGDLKTQIDQVEST	(SEQ ID NO: 22)
KTQIDQVESTAGSLQ	(SEQ ID NO: 23)
QVESTAGSLQGQWRG	(SEQ ID NO: 24)
AGSLQGQWRGAAGTA	(SEQ ID NO: 25)
GQWRGAAGTAAQAAV	(SEQ ID NO: 26)
AAGTAAQAAVVRFQE	(SEQ ID NO: 27)
AQAAVVRFQEAANKQ	(SEQ ID NO: 28)
VRFQEAANKQKQELD	(SEQ ID NO: 29)
AANKQKQELDEISTN	(SEQ ID NO: 30)
KQELDEISTNIRQAG	(SEQ ID NO: 31)
EISTNIRQAGVQYSR	(SEQ ID NO: 32)
IRQAGVQYSRADEEQ	(SEQ ID NO: 33)
VQYSRADEEQQQALS	(SEQ ID NO: 34)
ADEEQQQALSSQMGF	(SEQ ID NO: 35)

Page 21, replace the list starting at line 30 with:

Sequence of ESAT-6 (SEQ ID NO:_36):

MTEQQWNFAGIEAAASAIQGNVTSIHSLLDEGKQSLTKLAA AWGGSGSEAYQGVQQKWDATATELNNALQNLARTISEAGQ AMASTE GNVTGMFA

Page 22, replace the list starting at line 3 with:

Sequence of CFP-10 (SEQ ID NO:_37):

MAEMKTDAATLAQEAGNFERISGDLKTQIDQVESTAGSLQG QWRGAAGTAAQAAVVRFQEAANKQKQELDEISTNIRQAGV QYSRADEEQQQALS SQMGF